

Amendments to the Specification:

Please replace the paragraph at page 15, line 22, through page 16, line 3, with the following paragraph:

-- The screening method according to [24], which comprises measuring and comparing binding amount of the labeled peptide according to [1], its amide or ester, or salts thereof to cells comprising FPRL1, between (i) the case where (a) the labeled peptide according to [1], its amide or ester, or salts thereof, or (b) a labeled compound or a salt thereof that alters binding property between FPRL1 or a salt thereof and the peptide according to [1], its amide or ester, or salts thereof, is contacted with the cells comprising FPRL1, and (ii) the case where (a) the labeled peptide according to [1], its amide or ester, or salts thereof, or (b) a labeled compound or a salt thereof that alters binding property between FPRL1 or a salt thereof and the peptide according to [1], its amide or ester, or salts thereof, and a test compound are contacted with the cells comprising FPRL1;--

Please replace the paragraph at page 22, from line 7 through line 11, with the following paragraph:

--FIG. 4[[A]] shows a mass spectrum of porcine intrinsic FPRL1 ligand P3 by electrospray ionization mass spectrometer in which a mixture of polyvalent ions, trivalent ion m/z 575.30, 580.64 is shown. Horizontal axis and vertical axis represent mass/electric charge (m/z) and relative intensity in the case where the highest signal is referred to as 100%, respectively. In the figure, each of the numeric values associated with the labeled peaks indicates a value of m/z. --